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Reviewer Comments:
<210> 149
<211> 15
<212> DNA
<213> Artificial Sequence

<220>
<223> presence of
      the putative CodY box in yjcL

<220>
<221> misc_feature
<222> (1)..(15)

<400> 149
attgttagaa aatta   15

Regarding the above <220>-<222> section: what information pertains to
locations 1-15? Please indicate it in a <223> line. Same in Sequence
150.

<210> 179
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> presence of the putative CodY box in arcA

<<400> 179

atattttttg aaaaattttt taaaaa   26

Please remove the extra opening bracket "<" in the above <400> numeric
identifier: should just show <400>

<210> 216
<211> 26
<212> DNA
<213> Artificial Sequence

<220>
<223> WT

<400> 216
aatgttcaga aaattcatga acatac   26

As an explanation of "<213> Artificial Sequence", the above <223>
response needs more information regarding the source of the genetic
material.
****

Application No: 10562601   Version No: 2.0

Input Set:

Output Set:

Started: 2010-06-15 12:34:06.438

Finished: null

Elapsed: null

Total Warnings: 159

Total Errors: 1

No. of SeqIDs Defined: 234

Actual SeqID Count: 179

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W 213	Artificial or Unknown found in <213> in SEQ ID (2)
W 213	Artificial or Unknown found in <213> in SEQ ID (3)
W 213	Artificial or Unknown found in <213> in SEQ ID (4)
W 213	Artificial or Unknown found in <213> in SEQ ID (5)
W 213	Artificial or Unknown found in <213> in SEQ ID (6)
W 213	Artificial or Unknown found in <213> in SEQ ID (7)
W 213	Artificial or Unknown found in <213> in SEQ ID (8)
W 213	Artificial or Unknown found in <213> in SEQ ID (9)
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W 213	Artificial or Unknown found in <213> in SEQ ID (11)
W 213	Artificial or Unknown found in <213> in SEQ ID (12)
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W 213	Artificial or Unknown found in <213> in SEQ ID (14)
W 213	Artificial or Unknown found in <213> in SEQ ID (15)
W 402	Undefined organism found in <213> in SEQ ID (18)
W 213	Artificial or Unknown found in <213> in SEQ ID (24)
W 213	Artificial or Unknown found in <213> in SEQ ID (29)
W 213	Artificial or Unknown found in <213> in SEQ ID (39)
W 213	Artificial or Unknown found in <213> in SEQ ID (40)

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         Input Set:


         Output Set:



           Started:   2010-06-15 12:34:06.438
          Finished:   null
           Elapsed:   null
    Total Warnings:   159
      Total Errors:   1
No. of SeqIDs Defined: 234
 Actual SeqID Count:   179
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Error code	Error Description
W 213	Artificial or Unknown found in <213> in SEQ ID (41) This error has occured more than 20 times, will not be displayed
E 249	Order Sequence Error <223> -> <210>; Expected Mandatory Tag: <400> in SEQID (179)

<110> Hengst den, Christiaan D.

      Gajic, Olivera

      Kuipers, Oscar P.

      Kok, Jan

      Sikkema, Jan

      Geurts, Johannes M.W.

      Nauta, Arjen

<120> Methods and means for regulating gene expression

<130> P63590US00

<140> 10562601
<141> 2010-06-15

<150> PCT/NL2004/000474
<151> 2004-07-02

<150> EP 03077074.7
<151> 2003-07-02

<160> 234

<170> PatentIn Ver. 3.3

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<223> primer opp3

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<223> primer opp4

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<210> 8

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<210> 9
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<220>
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<212> DNA
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<220>
<223> primer HC-6

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<210> 14
<211> 15
<212> DNA
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<220>
<223> inversely repeated (IR) cis-element

<400> 14
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<220>
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<222> (2)..(2)
<223> Gly or Arg

<220>
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<222> (3)..(3)
<223> Met, Gly, Ile, Lys, Gln

<220>
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<222> (4)..(4)
<223> Arg or Gly

<220>
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<222> (5)..(5)
<223> Leu or Thr

<220>
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<222> (7)..(7)
<223> Thr or Ser

<400> 15
Gly Xaa Xaa Xaa Xaa Gly Xaa Thr
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<210> 16
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<212> PRT
<213> Escherichia coli

<220>
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<222> (1)..(8)
<223> /note="Putative GTP binding motif G1"

<400> 16
Leu Gly Gly Gly Thr Gly Thr Gly
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<210> 17
<211> 8
<212> PRT
<213> Bacillus subtilis

<220>
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<223> /note="Putative GTP binding motif G1"

<400> 17
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<210> 18
<211> 8
<212> PRT
<213> Bacillus halodurans

<220>
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<223> /note="Putative GTP binding motif G1"

<400> 18
Gly Gly Gln Arg Leu Gly Thr Leu
  1   5

<210> 19
<211> 8
<212> PRT

<213> Clostridium difficile

<220>
<221> BINDING
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<223> /note="Putative GTP binding motif G1"

<400> 19
Gly Gly Met Arg Leu Gly Ser Leu
 1   5

<210> 20
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<212> PRT
<213> Clostridium acetobutylicum

<220>
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<400> 20
Asn Arg Glu Arg Leu Gly Thr Leu
 1   5

<210> 21
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<212> PRT
<213> Streptococcus pneumoniae

<220>
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<400> 21
Ser Gly Ile Arg Leu Gly Ser Leu
 1   5

<210> 22
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<212> PRT
<213> Enterococcus faecalis

<220>
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<223> /note="Putative GTP binding motif G1"

<400> 22
Ala Gly Lys Arg Leu Gly Thr Ile
 1   5

<210> 23
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<220>
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<400> 23
Ser Gly Met Arg Leu Gly Thr Phe
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<210> 24
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<212> PRT
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<220>
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<220>
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<222> (2)..(2)
<223> Arg, Ala, Lys

<220>
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<222> (3)..(3)
<223> Val, Phe, Ile

<400> 24
Asp Xaa Xaa Gly
  1

<210> 25
<211> 4
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<220>
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<223> /note="Putative GTP binding motif G3"

<400> 25
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<210> 26

<211> 4
<212> PRT
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<220>
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Asp Arg Val Gly
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<400> 27
Asp Arg Ile Gly
  1

<210> 28
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<220>
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Asp Lys Ile Gly
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<210> 29
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<220>
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<220>
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<223> /note="Putative GTP binding motif G4"

<220>
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<223> Gly, Leu, Phe, Asn, Ser, Gln

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<210> 30
<211> 4
<212> PRT
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<220>
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<210> 31
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<210> 32
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<400> 33
Asn Glu Gly Ile
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<220>
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<400> 34
Ile Leu Asn Asp
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<210> 35
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<212> PRT
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<220>
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<223> /note="Putative GTP binding motif G4"

<400> 35
Leu Ile Ser Asp
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<210> 36
<211> 4
<212> PRT
<213> Enterococcus faecalis

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<223> /note="Putative GTP binding motif G4"

<400> 36
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<210> 37
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<400> 37
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<210> 38
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<400> 38
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<210> 39
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<210> 41
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<400> 43
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<210> 45
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<210> 49
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<400> 50
gaatttactg acgaatctat cattaa   26

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<400> 52
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<210> 66
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<220>
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<220>
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<400> 68
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<210> 69
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<210> 70
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<220>
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<210> 71
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<220>
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<210> 72
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<210> 73
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<220>
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<210> 74
<211> 15

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